



Additional file 8: Figure S1. Rarefaction curves of the sequencing reads. Each curve represents relationship between the number of sequencing reads and the number of observed OTUs (left, Hexapoda read data; right, Araneae read data). **a, e** Blocking primer A condition. **b, f** Blocking primer B condition. **c, g** Blocking primers A & B condition. **d, h** No blocking primer condition.